RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/5/6, 314
Source:	PCT
Date Processed by STIC:	01/05/2006

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PCT

RAW SEQUENCE LISTING DATE: 01/05/2006
PATENT APPLICATION: US/10/516,314 TIME: 11:29:36

Input Set : A:\P26360.ST5.txt

Output Set: N:\CRF4\01052006\J516314.raw

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3 <110> APPLICANT: Miyawaki, Atsushi
       Karasawa, Satoshi
6 <120> TITLE OF INVENTION: Chromoprotein
8 <130> FILE REFERENCE: P26360
10 <140> CURRENT APPLICATION NUMBER: 10/516,314
11 <141> CURRENT FILING DATE: 2004-12-10
13 <150> PRIOR APPLICATION NUMBER: PCT/JP03/07337
14 <151> PRIOR FILING DATE: 2003-06-10
16 <150> PRIOR APPLICATION NUMBER: JP2002-168584
17 <151> PRIOR FILING DATE: 2002-06-10
19 <160> NUMBER OF SEQ ID NOS: 12
21 <170> SOFTWARE: PatentIn version 3.3
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 229
25 <212> TYPE: PRT
26 <213> ORGANISM: Cnidopus japonicum
28 <400> SEQUENCE: 1
30 Met Ala Ser Met Ile Lys Asp Val Met Arg Phe Lys Met Asn Met Glu
34 Gly Thr Val Asn Gly His His Phe Lys Cys Glu Ala Asp Gly Glu Gly
38 Lys Pro Tyr Glu Gly Thr Gln Leu Gly Arg Ile Arg Val Thr Glu Gly
42 Gly Pro Leu Pro Phe Ser Phe Asp Ile Leu Ser Thr Cys Thr Gln Tyr
46 Gly Ser Lys Thr Met Ile Lys His Leu Ser Gly Ile Pro Asp Tyr Phe
                      70
                                           75
50 Lys Gln Ser Phe Pro Glu Gly Phe Thr Trp Glu Arg Asn Thr Ile Tyr
                                       90
54 Glu Asp Gly Gly His Leu Thr Ala His Gln Asp Thr Ser Leu Lys Gly
              100
                                   105
58 Asp Cys Leu Ile Tyr Lys Val Lys Val Leu Gly Gly Asn Phe Pro Ala
                               120
62 Asn Gly Pro Val Met Gln Lys Lys Ser Lys Gly Trp Glu Pro Cys Thr
                           135
                                               140
66 Glu Met Leu Tyr Pro Arg Asp Gly Val Leu Cys Gly Gln Thr Leu Met
                       150
                                           155
70 Ala Leu Lys Cys Ala Asn Gly Lys Asn Leu Thr Cys Gln Leu Arg Ser
                                       170
74 Thr Tyr Arg Ser Arg Lys Gln Ala Ser Ala Leu Gln Thr Pro Gly Phe
              180
                                   185
78 His Phe Gly Asp His Arg Ile Glu Ile Leu Lys Glu Ala Glu Gly Gly
          195
                               200
```

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```
82 Asn Tyr Phe Glu Gln His Glu Thr Ser Val Ala Arg Tyr Cys Asp Val
            210
                                215
                                                     220
     86 Ala Pro Ser Lys His
     87 225
     90 <210> SEQ ID NO: 2
     91 <211> LENGTH: 690
     92 <212> TYPE: DNA
     93 <213> ORGANISM: Cnidopus japonicus
     95 <400> SEQUENCE: 2
     96 atggcttcaa tgattaaaga cgttatgcgc ttcaagatga acatggaagg gacggttaac
                                                                               60
     98 ggccaccact tcaagtgtga agctgatgga gagggcaagc catatgaagg gactcaactc
                                                                              120
     100 ggaagaataa gggtcaccga aggcgggcca ttgccgtttt ctttcgacat cttgtcgact
                                                                               180
     102 tqcactcaat atqqaaqcaa qaccatqatc aaqcatttqt ccqqqattcc aqactacttt
                                                                               240
     104 aagcagtett ttecagaagg atteacetgg gaaaggaaca caatetatga ggatggtggt
     106 cacctcacag ctcaccaaga cactagtctt aagggagatt gcctgatcta taaagtgaaa
                                                                               360
     108 gtccttggag gtaattttcc tgccaatggc cctgtgatgc agaagaagag caaaggatgg
                                                                               420
     110 gaaccctgta ccgaaatgct ttatccacgt gatggagtgc tttgtggcca aacattgatg
                                                                                480
     112 gcacttaaat gcgccaatgg taaaaatctg acttgccagc taagatctac ttacaggtcc
                                                                               540
     114 agaaaacaag ccagtgcatt gcagacacca ggcttccatt tcggagacca tcgtattgag
                                                                               600
     116 atactcaagg aagcagagg gggcaattac tttgagcagc acgaaacatc agtcgccagg
                                                                                660
     118 tactgtgatg ttgccccgtc aaagcactga
                                                                                690
     121 <210> SEQ ID NO: 3
     122 <211> LENGTH: 21
     123 <212> TYPE: DNA
     124 <213> ORGANISM: Artificial sequence
     126 <220> FEATURE:
     127 <223> OTHER INFORMATION: Primer
     129 <400> SEQUENCE: 3
                                                                                 21
     130 gaaggrtgyg tcaayggrca y
     133 <210> SEQ ID NO: 4
     134 <211> LENGTH: 23
     135 <212> TYPE: DNA
     136 <213> ORGANISM: Artificial sequence
     138 <220> FEATURE:
     139 <223> OTHER INFORMATION: Primer
     141 <400> SEQUENCE: 4
     142 acvggdccat ydgvaagaaa rtt
                                                                                 23
     145 <210> SEQ ID NO: 5
     146 <211> LENGTH: 36
     147 <212> TYPE: DNA
     148 <213> ORGANISM: Artificial sequence
     150 <220> FEATURE:
     151 <223> OTHER INFORMATION: Primer
     154 <220> FEATURE:
     155 <221> NAME/KEY: misc feature
     156 <222> LOCATION: (24)..(35)
     157 <223> OTHER INFORMATION: wherein "n" is inosine
     159 <400> SEQUENCE: 5
W--> 160 ggccacgcgt cgactagtac gggnngggnn gggnng
                                                                                 36
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Input Set : A:\P26360.ST5.txt

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164 <211> LENGTH: 19
165 <212> TYPE: DNA
166 <213> ORGANISM: Artificial sequence
168 <220> FEATURE:
169 <223> OTHER INFORMATION: Primer
171 <400> SEQUENCE: 6
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172 atagatcagg caatctccc
175 <210> SEQ ID NO: 7
176 <211> LENGTH: 20
177 <212> TYPE: DNA
178 <213> ORGANISM: Artificial sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: Primer
183 <400> SEQUENCE: 7
                                                                            20
184 ggccacgcgt cgactagtac
187 <210> SEQ ID NO: 8
188 <211> LENGTH: 20
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial sequence
192 <220> FEATURE:
193 <223> OTHER INFORMATION: Primer
195 <400> SEQUENCE: 8
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196 ggtgagctgt gaggtgacca
199 <210> SEQ ID NO: 9
200 <211> LENGTH: 20
201 <212> TYPE: DNA
202 <213> ORGANISM: Artificial sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: Primer
207 <400> SEQUENCE: 9
                                                                            20
208 caccgaaggc gggccattgc
211 <210> SEQ ID NO: 10
212 <211> LENGTH: 44
213 <212> TYPE: DNA
214 <213 > ORGANISM: Artificial sequence
216 <220> FEATURE:
217 <223> OTHER INFORMATION: Primer
219 <400> SEQUENCE: 10
220 aactggaaga attcgcggcc gcagaatttt ttttttttt tttt
                                                                            44
223 <210> SEQ ID NO: 11
224 <211> LENGTH: 31
225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial sequence
228 <220> FEATURE:
229 <223> OTHER INFORMATION: Primer
231 <400> SEQUENCE: 11
232 cgggatccga ccatggcttc aatgattaaa g
                                                                            31
235 <210> SEQ ID NO: 12
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RAW SEQUENCE LISTING

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236 <211> LENGTH: 33

237 <212> TYPE: DNA

238 <213> ORGANISM: Artificial sequence

240 <220> FEATURE:

241 <223> OTHER INFORMATION: Primer

243 <400> SEQUENCE: 12

244 ccggaattct tagtgctttg acggggcaac atc

33

RAW SEQUENCE LISTING ERROR SUMMARY

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 24,25,29,30,34,35

VERIFICATION SUMMARY

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Input Set : A:\P26360.ST5.txt

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L:160 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0